

SEQ ID NO:1

SEQ ID NO:2

1 NCNNTNCCAT GTCGGCCCCA GTCACGNCAT ACTGANCATC TGACCGGGAT
51 ATAGTGTGGG TCCACACATC AGTCCCGACA CNAATGTGAT GTGGCACATA
101 AGGATTCTCC GCATANACAC AGCGACAATC TCGTCNGCAT AGTGGTAGGT
151 ATGATCNACA TGGGCCCCGAT CCATCTAACG GCGCACGCGG GACCACTTGT
201 CCTNATAGGT AATGCCCTGG CTACATGCTA CTTCTTTACT GTNCCCCCAC
251 CCCANCTACA CCGACNTNTT TNCCGGTCTA NATACACTCA ATATGCTGCC
301 CCTGCCCTCA TCGAACNGTC TGCACTNATA TACTGCAN